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1/1
GCC CTC GAG GGT ACC GAA TTC GCG CGT TCG GAA GGT GCT TCG GCG CTG GCG ACC GAT AAC
A L E G T E F A R S E G A S A L A T I N
61/21
CCG CTG AAG ACC ACT GTT GAA GAG TCG CTG TCG CGT GGA ATT GCT GGT AGC AAA ATT AAA
P L K T T V E E S L S R G I A G S K I K
121/41
ATT GGT ACT ACT GCT TCT ACT GCG ACC GAA ACA TAT GCC GGC GTC GAG CCG GAT GCC AAC
I G T T A S T A T E T Y A G V E P D A N
181/61
AAG TTG GGT GTA ATT GCT GTA GCA ATC GAA GAT AGT GGT GCG GGT GAT ATT ACC TTT ACC
K L G V I A V A I E D S G A G D I T F T
241/81
TTC CAG ACT GGT ACC TCT AGT CCC AAG AAT GCT ACT AAA GTT ATC ACT CTG AAC CGT ACT
F Q T G T S S P K N A T K V I T L N R T
301/101
GCC GAT GGG GTC TGG GCT TGT AAA TCT ACC CAG GAT CCG ATG TTC ACT CCG AAA GGT TTT
A D G V W A C K S T Q D P M F T P K G S
361/121
GAT AAC
D N

Fig. 1C



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DNA sequence 610 b.p. ttctagataacg ... CGAAAGCTTGGG linear

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86/1      116/11
gcg ctc gag cac cat cat cac cat ggt ggt ggt ggc gag att gag gcc ctc aag gct gaa
A L E H H H H H G G G G E I E A L K A E
146/21      176/31
atc gaa gcc cta aag gcc gag ata gaa gca ctt aag gca gag atc gag gcg cta aaa gcg
I E A L K A E I E A L K A E I E A L K A
206/41      236/51
gaa ata gag gct ctg aag gca ggc ggt gga gga gaa ttc GCG CGT TCG GAA GGT GCT TCG
E I E A L K A G G G G E F A R S E G A S
266/61      296/71
GCG CTG GCG ACG ATC AAC CCG CTG AAG ACC ACT GTT GAA GAG TCG CTG TCG CGT GGA ATT
A L A T I N P L K T T V E E S L S R G I
326/81      356/91
GCT GGT AGC AAA ATT AAA ATT GGT ACT ACT GCT TCT ACT GCG ACC GAA ACA TAT GCC GGC
A G S K I K I G T T A S T A T E T Y A G
386/101      416/111
GTC GAG CCG GAT GCC AAC AAG TTG GGT GTA ATT GCT GTA GCA ATC GAA GAT AGT GGT GCG
V E P D A N K L G V I A V A I E D S G A
446/121      476/131
GGT GAT ATT ACC TTT ACC TTC CAG ACT GGT ACC TCT AGT CCC AAG AAT GCT ACT AAA GTT
G D I T F T F Q T G T S S P K N A T K V
506/141      536/151
ATC ACT CTG AAC CGT ACT GCG GAT GGG GTC TGG GCT TGT AAA TCT ACC CAG GAT CCG ATG
I T L N R T A D G V W A C K S T Q D P M
566/161
TTC ACT CCG AAA GGT TGT GAT AAC
F T P K G G C D N
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Fig. 5A

DNA sequence 610 b.p. ttctagataacg ... CGAAAGCTTGGG linear

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86/1      116/11
gcg ctc gag cac cat cat cac cat ggt ggt ggt ggc gag gta tcc gct tta gag aaa gaa
A L E H H H H H G G G G E V S A L E K E
146/21      176/31
gtt tct gct ctc gaa aaa gag gtc agt gct ctg gaa aaa gag gtc tca gcc ttg gaa aag
V S A L E K E V S A L E K E V S A L E K
206/41      236/51
gaa gta tca gca ctt gag aag ggc ggt gga gga gaa ttc GCG CGT TCG GAA GGT GCT TCG
E V S A L E K G G G G E F A R S E G A S
266/61      296/71
GCG CTG GCG ACG ATC AAC CCG CTG AAG ACC ACT GTT GAA GAG TCG CTG TCG CGT GGA ATT
A L A T I N P L K T T V E E S L S R G I
326/81      356/91
GCT GGT AGC AAA ATT AAA ATT GGT ACT ACT GCT TCT ACT GCG ACC GAA ACA TAT GCC GGC
A G S K I K I G T T A S T A T E T Y A G
386/101      416/111
GTC GAG CCG GAT GCC AAC AAG TTG GGT GTA ATT GCT GTA GCA ATC CAA GAT ACT GCT GCG
V E P D A N K L G V I A V A I E D S G A
446/121      476/131
GGT GAT ATT ACC TTT ACC TTC CAG ACT GGT ACC TCT AGT CCC AAG AAT GCT ACT AAA GTT
G D I T F T F Q T G T S S P K N A T K V
506/141      536/151
ATC ACT CTG AAC CGT ACT GCG GAT GGG GTC TGG GCT TGT AAA TCT ACC CAG GAT CCG ATG
I T L N R T A D G V W A C K S T Q D P M
566/161
TTC ACT CCG AAA GGT TGT GAT AAC
F T P K G G C D N
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Fig. 5B